

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/518, 297A
Source:	1FW16
Date Processed by STIC:	1/24/05
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

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- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
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 Alexandria, VA 22314

Revised 01/24/05



IFW16

RAW SEQUENCE LISTING DATE: 01/24/2005
PATENT APPLICATION: US/09/518,297A TIME: 11:17:44

Input Set : A:\54600-8130US00.TXT

Output Set: N:\CRF4\01242005\I518297A.raw

```
4 <110> APPLICANT: Lim, Moon Young
       Edwards, Cynthia A.
 6
        Fry, Kirk E.
 7
        Bruice, Thomas W.
        Starr, Douglas B.
9
        Laurance, Megan E.
10
        Kwok, Yan
13 <120> TITLE OF INVENTION: DNA Binding Compound-Mediated Molecular
       Switch System
16 <130> FILE REFERENCE: 54600-8130US00
18 <140> CURRENT APPLICATION NUMBER: US 09/518,297A
19 <141> CURRENT FILING DATE: 2000-03-03
21 <150> PRIOR APPLICATION NUMBER: US 60/122,513
22 <151> PRIOR FILING DATE: 1999-03-03
24 <150> PRIOR APPLICATION NUMBER: US 60/154,605
                                                                 Does Not Comply
25 <151> PRIOR FILING DATE: 1999-09-17
27 <160> NUMBER OF SEQ ID NOS: 84
                                                             Corrected Diskette Needer
29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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ERRORED SEQUENCES

misaligned amino acid rumbering

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VERIFICATION SUMMARY

DATE: 01/24/2005 TIME: 11:17:46

PATENT APPLICATION: US/09/518,297A

Input Set : A:\54600-8130US00.TXT

Output Set: N:\CRF4\01242005\I518297A.raw

L:61 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:65 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:66 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:180 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:184 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:188 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:192 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:223 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:227 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:1255 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:64
M:332 Repeated in SeqNo=64
L:1257 M:252 E: No. of Seq. differs, <211> LENGTH:Input:43 Found:44 SEQ:64
L:1382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75 after pos.:0
L:1391 M:283 W: Missing Blank Line separator, <400> field identifier

L:1470 M:283 W: Missing Blank Line separator, <400> field identifier